

CLAIMS

WHAT IS CLAIMED IS:

1. An isolated nucleic acid comprising at least 10 contiguous nucleotides of a sequence selected from the group consisting of the human polynucleotide coding sequences hRxx-yyy shown in Tables 1-129, or its complement.
2. A host cell comprising a recombinant nucleic acid of claim 1.
3. An expression vector comprising the isolated nucleic acid according to claim 1.
4. A host cell comprising the expression vector of claim 3.
5. The polynucleotide according to claim 1, wherein said polynucleotide, or its complement or a fragment thereof, further comprises a detectable label.
6. The polynucleotide according to claim 1, wherein said polynucleotide, or its complement or a fragment thereof, is attached to a solid support.
7. The polynucleotide according to claim 1, wherein said polynucleotide, or its complement or a fragment thereof, is prepared at least in part by chemical synthesis.
8. The polynucleotide according to claim 1, wherein said polynucleotide, or its complement or a fragment thereof, is an antisense fragment.
9. The polynucleotide according to claim 1, wherein said polynucleotide, or its complement or a fragment thereof, is single stranded.
10. The polynucleotide according to claim 1, wherein said polynucleotide, or its complement or a fragment thereof, is double stranded.
11. The polynucleotide according to claim 1, comprising at least 15 contiguous nucleotides.
12. The polynucleotide according to claim 1, comprising at least 20 contiguous nucleotides.
13. A microarray for detecting a cancer associated (CA) nucleic acid comprising:
at least one probe comprising at least 10 contiguous nucleotides of a sequence selected from the group consisting of the human polynucleotide coding sequences hRxx-yyy shown in Tables 1-129, or its complement.

14. The microarray according to claim 13, comprising at least 15 contiguous nucleotides.
15. The microarray according to claim 13, comprising at least 20 contiguous nucleotides.
16. An isolated polypeptide, encoded within an open reading frame of a CA sequence selected from the group consisting of the human genomic polynucleotide sequences hDxx-yyy shown in Tables 1-129, or its complement.
17. The polypeptide of claim 16, wherein said polypeptide comprises the amino acid sequence encoded by a polynucleotide selected from the group consisting of hRxx-yyy shown in Tables 1-129.
18. The polypeptide of claim 16, wherein said polypeptide comprises the amino acid sequence encoded by a polypeptide selected from the group consisting of hPxx-yyy shown in Tables 1-129.
19. The polypeptide of claim 16, wherein said polypeptide comprises the amino acid sequence of an epitope of the amino acid sequence of a CA polypeptide selected from the group consisting of hPxx-yyy shown in Tables 1-129.
20. The polypeptide of claim 16, wherein said polypeptide or fragment thereof is attached to a solid support.
21. An isolated antibody or antigen binding fragment thereof, that binds to a polypeptide according to anyone of claims 16-20.
22. The isolated antibody or antigen binding fragment thereof according the claim 21, wherein said antibody or fragment thereof is attached to a solid support.
23. The isolated antibody or antigen binding fragment thereof according the claim 21, wherein said antibody is a monoclonal antibody.
24. The isolated antibody or antigen binding fragment thereof according the claim 21, wherein said antibody is a polyclonal antibody.
25. The isolated antibody or antigen binding fragment thereof according the claim 21, wherein said antibody or fragment thereof further comprises a detectable label.
26. An isolated antibody that binds to a polypeptide, or antigen binding fragment thereof, according to any of claims 16-20, prepared by a method comprising the following

steps of: (i) immunizing a host animal with a composition comprising said polypeptide, or antigen binding fragment thereof, and ii) collecting cells from said host expressing antibodies against the antigen or antigen binding fragment thereof.

27. A kit for diagnosing the presence of cancer in a test sample, said kit comprising at least one polynucleotide that selectively hybridizes to a CA polynucleotide sequence selected from the group consisting of the polynucleotide sequences hDxx-yyy shown in Tables 1-129, a fragment thereof, or their complement.

28. A kit for diagnosing the presence of cancer in a test sample, said kit comprising at least one polynucleotide that selectively hybridizes to the sequence of a polynucleotide sequence selected from the group consisting of the polynucleotide sequences hRxx-yyy shown in Tables 1-129, a fragment thereof, or their complement.

29. An electronic library comprising a polynucleotide, or fragment thereof, comprising a CA polynucleotide sequence selected from the group consisting of the polynucleotide sequences hDxx-yyy shown in Tables 1-129.

30. An electronic library comprising a polynucleotide, or fragment thereof, comprising a CA polynucleotide sequence selected from the group consisting of the polynucleotide sequences hRxx-yyy shown in Tables 1-129.

31. An electronic library comprising a polypeptide, or fragment thereof, comprising a CA polypeptide sequence selected from the group consisting of the polynucleotide sequences hPxx-yyy shown in Tables 1-129.

32. A method for screening for anticancer activity in a potential drug, the method comprising:

(a) providing a cell that expresses a cancer associated (CA) gene encoded by a nucleic acid sequence selected from the group consisting of the sequences hDxx-yyy shown in Tables 1-129 or fragment thereof;

(b) contacting a tissue sample derived from a cancer cell with an anticancer drug candidate; and

(c) monitoring an effect of the anticancer drug candidate on an expression of the CA gene in the tissue sample.

33. The method of screening for anticancer activity according to claim 32, wherein the CA gene comprises at least one nucleic acid sequence selected from the group consisting of the sequences hRxx-yyy shown in Tables 1-129.

34. The method of screening for anticancer activity according to claim 32, further comprising:

(d) comparing the level of expression of the in the absence of said drug candidate to the level of expression in the presence of the drug candidate.

35. The method of screening for anticancer activity according to claim 33, wherein the drug candidate is an inhibitor of transcription and modulates the activity of a CAP sequence (hPxx-yyy) selected from the group consisting of SEQ ID NOS: 28, 42, 56, 142, 167, 169, 173, 175, 177, 185, 187, 193, 195, 227, 255, 258, 260, 274, 286, 298, 300, 302, 304, 322, 324, 392, 394, 412, 460, 462, 636, 728, 789, 861, 901, 907, 927, and 977.

36. The method of screening for anticancer activity according to claim 33, wherein the drug candidate is a G-protein coupled receptor antagonist and modulates the activity of a CAP sequence (hPxx-yyy) selected from the group consisting of SEQ ID NOS: 46, 913, 915, 921, and 933.

37. The method of screening for anticancer activity according to claim 33, wherein the drug candidate is a calcium binding protein antagonist and modulates the activity of a CAP sequence (hPxx-yyy) selected from the group consisting of SEQ ID NOS: 48, 50, 201, 336, 338, 346, 404, and 406.

38. The method of screening for anticancer activity according to claim 33, wherein the drug candidate is a ubiquitin cycle antagonist and modulates the activity of a CAP sequence (hPxx-yyy) selected from the group consisting of SEQ ID NOS: 34, 36, 62, 104, 106, 108, 110, 151, and 153.

39. A method for detecting cancer associated with expression of a polypeptide in a test cell sample, comprising the steps of:

(i) detecting a level of expression of at least one polypeptide selected from the group consisting of hPxx-yyy shown in Tables 1-129, or a fragment thereof; and

(ii) comparing the level of expression of the polypeptide in the test sample with a level of expression of polypeptide in a normal cell sample, wherein an altered level of expression of the polypeptide in the test cell sample relative to the level of polypeptide

expression in the normal cell sample is indicative of the presence of cancer in the test cell sample.

40. A method for detecting cancer associated with expression of a polypeptide in a test cell sample, comprising the steps of:

(i) detecting a level of activity of at least one polypeptide selected from the group consisting of hPxx-yyy shown in Tables 1-129, or a fragment thereof, wherein said activity corresponds to at least one activity for the polypeptide listed in Table 130; and

(ii) comparing the level of activity of the polypeptide in the test sample with a level of activity of polypeptide in a normal cell sample, wherein an altered level of activity of the polypeptide in the test cell sample relative to the level of polypeptide activity in the normal cell sample is indicative of the presence of cancer in the test cell sample.

41. A method for detecting cancer associated with the presence of an antibody in a test serum sample, comprising the steps of:

(i) detecting a level of an antibody against an antigenic polypeptide selected from the group consisting of hPxx-yyy shown in Tables 1-129, or antigenic fragment thereof; and

(ii) comparing said level of said antibody in the test sample with a level of said antibody in the control sample, wherein an altered level of antibody in said test sample relative to the level of antibody in the control sample is indicative of the presence of cancer in the test serum sample.

42. A method for screening for a bioactive agent capable of modulating the activity of a CA protein (CAP), wherein said CAP is encoded by a nucleic acid comprising a nucleic acid sequence selected from the group consisting of the polynucleotide sequences hRxx-yyy shown in Tables 1-129, said method comprising:

a) combining said CAP and a candidate bioactive agent; and

b) determining the effect of the candidate agent on the bioactivity of said CAP.

43. The method of screening for the bioactive agent according to claim 42, wherein the bioactive agent affects the expression of the CA protein (CAP).

44. The method of screening for the bioactive agent according to claim 42, wherein the bioactive agent affects the activity of the CA protein (CAP), wherein such activity is selected from the activities listed in Table 130.

45. The method of screening for the bioactive agent according to claim 42, wherein the bioactive agent is an inhibitor of transcription and modulates the activity of a CAP sequence (hPxx-yyy) selected from the group consisting of SEQ ID NOS: 28, 42, 56, 142, 167, 169, 173, 175, 177, 185, 187, 193, 195, 227, 255, 258, 260, 274, 286, 298, 300, 302, 304, 322, 324, 392, 394, 412, 460, 462, 636, 728, 789, 861, 901, 907, 927, and 977.

46. The method of screening for the bioactive agent according to claim 42, wherein the bioactive agent is a G-protein coupled receptor antagonist and modulates the activity of a CAP sequence (hPxx-yyy) selected from the group consisting of SEQ ID NOS: 46, 913, 915, 921, and 933.

47. The method of screening for the bioactive agent according to claim 42, wherein the bioactive agent is a calcium binding protein antagonist and modulates the activity of a CAP sequence (hPxx-yyy) selected from the group consisting of SEQ ID NOS: 48, 50, 201, 336, 338, 346, 404, and 406.

48. The method of screening for the bioactive agent according to claim 42, wherein the bioactive agent is a ubiquitin cycle antagonist and modulates the activity of a CAP sequence (hPxx-yyy) selected from the group consisting of SEQ ID NOS: 34, 36, 62, 104, 106, 108, 110, 151, and 153.

49. A method for diagnosing cancer comprising:

a) determining the expression of one or more genes comprising a nucleic acid sequence selected from the group consisting of the human sequences outlined in Tables 1-129, in a first tissue type of a first individual; and

b) comparing said expression of said gene(s) from a second normal tissue type from said first individual or a second unaffected individual;

wherein a difference in said expression indicates that the first individual has cancer.

50. A method for treating cancers comprising administering to a patient an inhibitor of a CA protein (CAP), wherein said CAP is encoded by a nucleic acid comprising a human nucleic acid sequence selected from the group consisting of the sequences outlined in Tables 1-129.

51. The method for treating cancers according to claim 50, wherein the inhibitor of a CA protein (CAP) binds to the CA protein.
52. The method for treating cancers according to claim 50, wherein the inhibitor of a CA protein (CAP) is an inhibitor of transcription and modulates the activity of a CAP sequence (hPxx-yyy) selected from the group consisting of SEQ ID NOS: 28, 42, 56, 142, 167, 169, 173, 175, 177, 185, 187, 193, 195, 227, 255, 258, 260, 274, 286, 298, 300, 302, 304, 322, 324, 392, 394, 412, 460, 462, 636, 728, 789, 861, 901, 907, 927, and 977.
53. The method for treating cancers according to claim 50, wherein the inhibitor of a CA protein (CAP) is a G-protein coupled receptor antagonist and modulates the activity of a CAP sequence (hPxx-yyy) selected from the group consisting of SEQ ID NOS: 46, 913, 915, 921, and 933.
54. The method for treating cancers according to claim 50, wherein the inhibitor of a CA protein (CAP) is a calcium binding protein antagonist and modulates the activity of a CAP sequence (hPxx-yyy) selected from the group consisting of SEQ ID NOS: 48, 50, 201, 336, 338, 346, 404, and 406.
55. The method for treating cancers according to claim 50, wherein the inhibitor of a CA protein (CAP) is a ubiquitin cycle antagonist and modulates the activity of a CAP sequence (hPxx-yyy) selected from the group consisting of SEQ ID NOS: 34, 36, 62, 104, 106, 108, 110, 151, and 153.